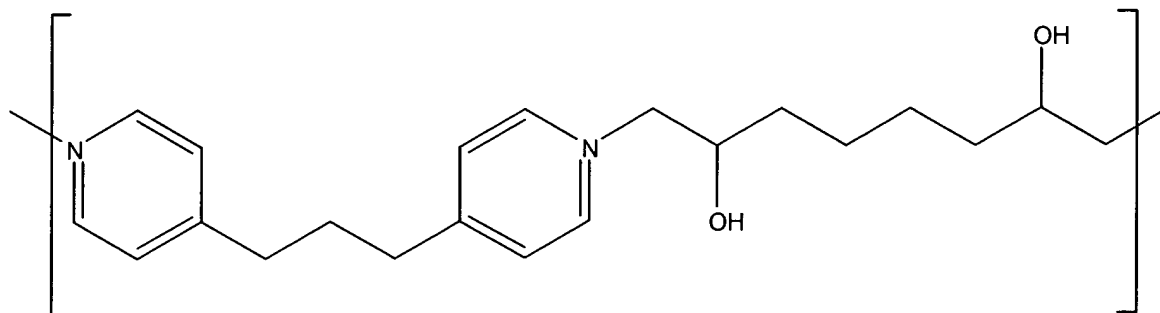


10/051,765

-2-



Claims readable on the elected species are 36-38, 42-44 and 48-50. Claims 39-41 and 45-47 (from Group VIII) also read on the elected polymer species.

Please amend the application as follows:

In the Specification

Please replace the table at page 30 with the following table:

TABLE I
Polymer MIC(μ g/ml)

| Formula # | S.aureus (ATCC 29213) | MRSA (5 strains) ⁽¹⁾ | GISA (5 strains) ⁽²⁾ | E. faecalis (ATCC 29212) | VRE (4 strains) | P. aeruginosa (ATCC 27853) | Candida* | B. cepacia | Oral pathogens ** | C. difficile | H. pylori (Win) | E. coli (ATCC 25922) | S. marcescens |
|-----------|-----------------------|---------------------------------|---------------------------------|--------------------------|-----------------|----------------------------|----------|------------|-------------------|--------------|-----------------|----------------------|---------------|
| VIII | 0.65 | 1-2 | 1 | 1 | 1-3 | 10.4 | 0.65-1.3 | 41.5 | nd | 0.65 | 0.2 | 1.3 | 3 |
| II | 0.31 | 1.9 | ≤ 0.125 | 0.31 | 0.3 | 1.3 | 1.3-2.6 | 5.2 | nd | <0.16 | 0.2 | .65 | 3 |
| IV | 0.65 | 2.2-3.8 | nd | 0.65 | 2-2.6 | 2.6 | 0.32-2.6 | 5.2 | nd | nd | 3 | 1.3 | 9 |
| II | 1.8 | 1-2.6 | nd | 1 | 1-2 | 9 | 3-28 | 28 | nd | nd | 1 | 1 | 3 |
| III | 1.4 | 0.4-1.8 | nd | 3 | 1-3 | 9 | 1-3 | 28 | nd | nd | 2 | 3 | 3 |
| XII | 0.31 | 0.3-0.5 | nd | 0.3 | 0.3 | 0.31 | 0.16-2.6 | 41.5 | nd | nd | nd | 2.6 | 3 |
| XIX | 0.65 | 1 | 0.5 | 1 | 1-3 | 0.65 | 1.3-2.6 | 41.5 | <10 | nd | 1 | 2.6 | 3 |
| XX | 1.3 | 2.2-3 | 16 | 1 | 1-3 | 1.3 | 2.6 | 5.2 | <10 | <0.16 | 0.4 | 1.3 | 3 |
| XXIII | 1 | 1.4-2.2 | nd | 1 | 1-3 | 9 | 3 | 9 | nd | nd | 1 | 3 | 3 |
| IX+ | 1 | nd | nd | 0.3 | nd | 83 | 1-9 | 28 | nd | nd | nd | 3 | 28 |
| IX++ | 1 | 1 | nd | 3 | 3-20 | 83 | 1-9 | 9 | nd | nd | 0.7 | 3 | 9 |
| IX+++ | 0.65 | nd | 8 | 1.25 | nd | 5.2 | 0.65-1.3 | 5.2 | <10 | 2.5 | 1.3 | 2.6 | 3 |

* 3 strains *C. glabrata*, 6 strains *C. albicans*, 2 strains *C. tropicalis*, 2 strains *C. Krusei*; ** including: *Actinobacillus actinomycetemcomitans*, *Porphyromonas gingivalis*, *Prevotella intermedia*, *Campylobacter rectus*, *Treponema denticola*, *Streptococcus intermedius*, *Fusobacterium nucleatum*, *Streptococcus gordonii*, and *Actinomyces viscosus*; ¹ methicillin resistant *Staphylococcus aureus*; ² glycopeptide intermediate *Staphylococcus aureus*; + (where R7 is butyl), ++(where R7 is decyl), +++(where R7 is dodecyl)